

kol447		(G, T, S, A) (I, L, M, F, P, W, Y, V) RR (I, L, M, F, P, W, Y, V) (I, L, M, F, P, W, Y, V) (G, T)
	1:	(G) (I) RR (F) (I) (G) (S) (I) (W) R (F) (I) R GIRRFIGSIWRFIR AFYG
ADO34356	ck: 8202 len: 14	! Add34356 Synthetic apolipoprotein-E mimicki
		(G, T, S, A) (I, L, M, F, P, W, Y, V) RR (I, L, M, F, P, W, Y, V) (I, L, M, F, P, W, Y, V) (G, T)
	1:	(G) (I) RR (F) (Y) (G) (S) (I) (W) R (F) (I) R GIRRFIGSIWRFIR
ADO34364	ck: 8241 len: 14	! Add34364 Synthetic apolipoprotein-E mimicki
		(G, T, S, A) (I, L, M, F, P, W, Y, V) RR (I, L, M, F, P, W, Y, V) (I, L, M, F, P, W, Y, V) (G, T)
	1:	(G) (I) RR (F) (Y) (G) (S) (I) (W) R (F) (I) R GIRRFIGSIWRFIR
ADO34419	ck: 8198 len: 14	! Add34419 Synthetic apolipoprotein-E mimicki
		(G, T, S, A) (I, L, M, F, P, W, Y, V) RR (I, L, M, F, P, W, Y, V) (I, L, M, F, P, W, Y, V) (G, T)
	1:	(G) (I) RR (F) (Y) (G) (S) (I) (W) R (F) (I) R GIRRFIGSIWRFIR
ADO34227	ck: 2959 len: 18	! Add34227 Synthetic apolipoprotein-E mimicki
		(G, T, S, A) (I, L, M, F, P, W, Y, V) RR (I, L, M, F, P, W, Y, V) (I, L, M, F, P, W, Y, V) (G, T)
	1:	(G) (I) RR (F) (Y) (G) (S) (I) (W) R (F) (I) R GIRRFIGSIWRFIR AFVG
ADO34344	ck: 2867 len: 18	! Add34344 Synthetic apolipoprotein-E mimicki
		(G, T, S, A) (I, L, M, F, P, W, Y, V) RR (I, L, M, F, P, W, Y, V) (I, L, M, F, P, W, Y, V) (G, T)
	1:	(G) (I) RR (F) (Y) (G) (S) (I) (W) R (F) (I) R GIRRFIGSIWRFIR AFVG
ADO34372	ck: 8211 len: 14	! Add34372 Synthetic apolipoprotein-E mimicki
		(G, T, S, A) (I, L, M, F, P, W, Y, V) RR (I, L, M, F, P, W, Y, V) (I, L, M, F, P, W, Y, V) (G, T)
	1:	(G) (I) RR (F) (Y) (G) (S) (I) (W) R (F) (I) R GFRRIYGSIWRFIR
ADO34333	ck: 3007 len: 18	! Add34333 Synthetic apolipoprotein-E mimicki
		(G, T, S, A) (I, L, M, F, P, W, Y, V) RR (I, L, M, F, P, W, Y, V) (I, L, M, F, P, W, Y, V) (G, T)
	1:	(G) (I) RR (F) (Y) (G) (S) (I) (W) R (F) (I) R GIRRIIGSFWRFFR AFYG
ADO34339	ck: 2998 len: 18	! Add34339 Synthetic apolipoprotein-E mimicki
		(G, T, S, A) (I, L, M, F, P, W, Y, V) RR (I, L, M, F, P, W, Y, V) (I, L, M, F, P, W, Y, V) (G, T)
	1:	(G) (I) RR (F) (Y) (G) (S) (I) (W) R (F) (I) R GLRRFIGSIWRFIR AFYG
ADO34233	ck: 3010 len: 18	! Add34233 Synthetic apolipoprotein-E mimicki
		(G, T, S, A) (I, L, M, F, P, W, Y, V) RR (I, L, M, F, P, W, Y, V) (I, L, M, F, P, W, Y, V) (G, T)
	1:	(G) (I) RR (F) (Y) (G) (S) (I) (W) R (F) (I) R GIRRFIGSIWRFIR AFYG
ADO34335	ck: 3019 len: 18	! Add34335 Synthetic apolipoprotein-E mimicki
		(G, T, S, A) (I, L, M, F, P, W, Y, V) RR (I, L, M, F, P, W, Y, V) (I, L, M, F, P, W, Y, V) (G, T)
	1:	(G) (I) RR (F) (Y) (G) (S) (I) (W) R (F) (I) R GFRRIYGSIWRFIR

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ADO34350 ck: 3064 len: 18 ! Ado34350 Synthetic apolipoprotein-E mimicki
(G,T,S,A)(I,L,M,F,P,W,Y,V)RR(I,L,M,F,P,W,Y,V)(I,L,M,F,P,W,Y,V)(G,T
(G)(L)RR(F)(I)(G)(S)(L)(W)R(F)(L)R
GLRRFIGSLWRFLR AFYG

ADO34378 ck: 8241 len: 14 ! Ado34378 Synthetic apolipoprotein-E mimicki
(G,T,S,A)(I,L,M,F,P,W,Y,V)RR(I,L,M,F,P,W,Y,V)(I,L,M,F,P,W,Y,V)(G,T
(G)(I)RR(F)(Y)(G)(S)(I)(W)R(F)(L)R
GIRRFYGSIWWRFLR

ADX96509 ck: 8299 len: 671 ! Adx96509 Plant full length insert polypepti
(G,T,S,A)(I,L,M,F,P,W,Y,V)RR(I,L,M,F,P,W,Y,V)(I,L,M,F,P,W,Y,V)(G,T
(A)(L)RR(L)(V)(A)(L)(P)R(L)(P)R
ALRRLVAAALPRLPR AGEVA

ADY07006 ck: 750 len: 646 ! Ady07006 Plant full length insert polypepti
(G,T,S,A)(I,L,M,F,P,W,Y,V)RR(I,L,M,F,P,W,Y,V)(I,L,M,F,P,W,Y,V)(G,T
(A)(L)RR(L)(V)(A)(A)(L)(P)R(L)(P)R
ALRRLVAAALPRLPR AGEVA

k01447 1: RRDHA 37: RRDHA

```

Databases searched:

EMBL, Release 8.0, Released on 4Apr2006, Formatted on 29Apr2006

```

Total finds: 45
Total length: 457,216,429
Total sequences: 2,589,679
CPU time: 32:07.06

```

```
! FINDPATTERNS on pir:* allowing 0 mismatches
! Using patterns from: kol447.dat  February 9, 2007 16:28 ..
.

Databases searched:
  NBRF, Release 80.0, Released on 31Dec2004, Formatted on 21Jun2005

Total finds: 0
Total length: 96,216,763
Total sequences: 283,416
CPU time: 05:41.52
```

No results from PIR.

```
! FINDPATTERNS on uniprot:* allowing 0 mismatches
! Using patterns from: kol447.dat  February 9, 2007 16:28 ..
Q6KA74_ORYSA ck: 7693 len: 633  ! Q6ka74 oryza sativa (japonica cultivar-group)
kol447          (G,T,S,A) (I,L,M,P,W,Y,V)RR(I,L,M,P,W,Y,V) (I,L,M,F,P,W,Y,V) (G,T
(S) (L) RR(L) (V) (A) (G) (L) (P) R(L) (P) R
23 : RRDHA SLRLVAGLPRLPR AGEVA
```

Databases searched:
UNIPROT, Release 7.2, Released on 7Mar2006, Formatted on 7Mar2006

Total finds:	1
Total length:	925,015,592
Total sequences:	2,843,598
CPU time:	56:34.48

Result from Uniprot